

Figure 1A

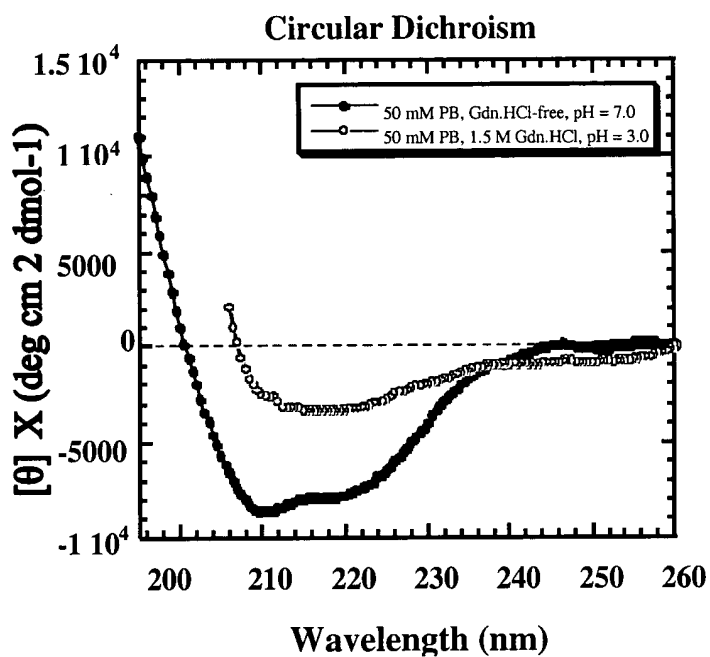
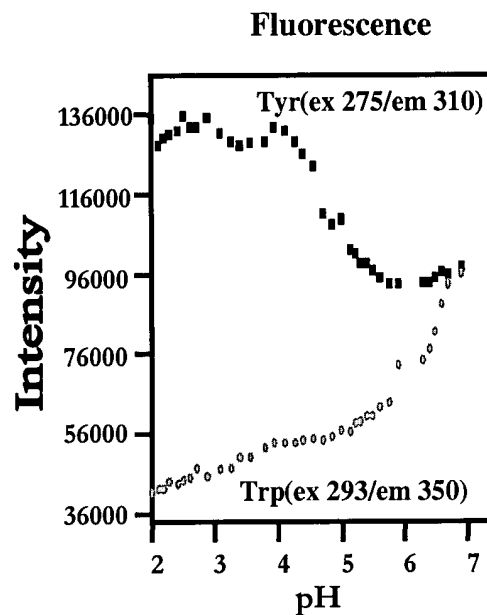


Figure 1B



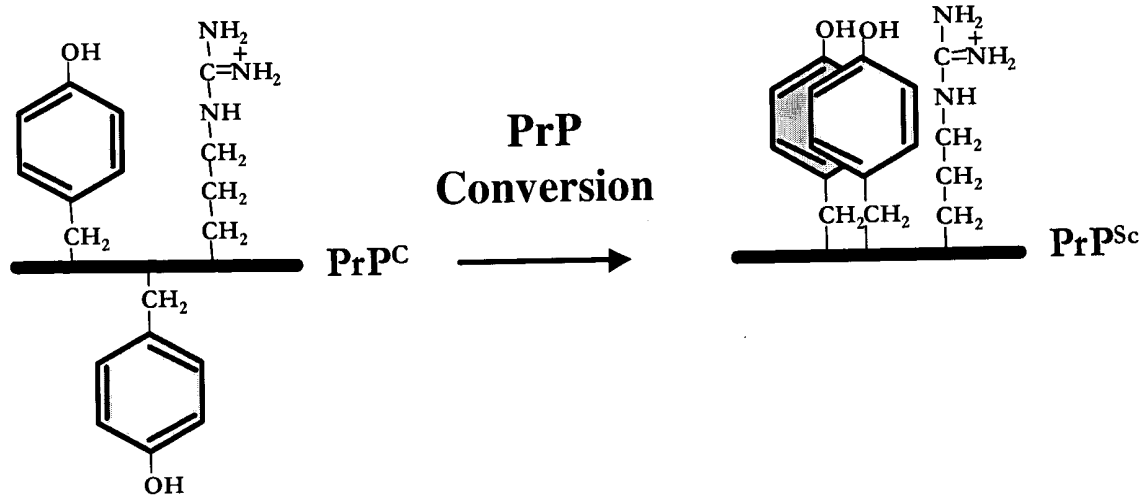
[illegible]

BOVINE	1	MVKSHIGSWILVLFVAMWS	DVGLCKKRPKPGGG	WNTGGS	SRYPGQGS	PGGNR	YPPQGGG	GW
MAN	1	- - MANLGSWMLVLFVATWS	DLGLCKKRPKPGG	- WNTGGS	SRYPGQGS	PGGNR	YPPQGGG	GW
SHEEP	1	MVKSHIGSWILVLFVAMWS	DVGLCKKRPKPGGG	WNTGGS	SRYPGQGS	PGGNR	YPPQGGG	GW
MOUSE	1	- - MANLGYWL LALFVT MWT	DVGLCKKRPKPGG	- WNTGGS	SRYPGQGS	PGGNR	YPPQGGG	- TW
HAMSTER	1	- - MANLSYWL LALFVAMWT	DVGLCKKRPKPGG	- WNTGGS	SRYPGQGS	PGGNR	YPPQGGG	TW
BOVINE	58	GQPHGGGGWG	QPHGGGGWG	QPHGGGGW	QPHGGG	- WG	QPHGGGGGGWG	QGG THGQWNKPSKPKTN
MAN	58	GQPHGGGGWG	QPHGGGGWG	QPHGGGGW	QPHGGG	- WG	Q- - GG	- - - - - THSQWNKPSKPKTN
SHEEP	61	GQPHGGGGWG	QPHGGGGWG	QPHGGGGW	QPHGGG	- GG	- - - - -	- THSQWNKPSKPKTN
MOUSE	57	GQPHGGGGWG	QPHGGGGWG	QPHGGGGW	QPHGGG	- WG	Q- - GG	- - - - - THNQWNKPSKPKTN
HAMSTER	58	GQPHGGGGWG	QPHGGGGWG	QPHGGGGW	QPHGGG	- WG	Q- - GG	- - - - - THNQWNKPSKPKTN
BOVINE	120	MKHVAGAAAAAGAVVGG	LGGYMLGS	AMSRPLIHFG	SDYEDR	YYR	REN	MHRRYPNQVYYR PVDQ
MAN	109	MKHMAGAAAAAGAVVGG	LGGYMLGS	AMSRPLIHFG	SDYEDR	YYR	REN	MHRRYPNQVYYR PVDQ
SHEEP	112	MKHVAGAAAAAGAVVGG	LGGYMLGS	AMSRPLIHFG	NDYEDR	YYR	REN	MYRRYPNQVYYR PVDQ
MOUSE	108	LKHVAGAAAAAGAVVGG	LGGYMLGS	AVSRPMIHFG	NDWE	DR	YYR	RENMYRRYPNQVYYR PVDQ
HAMSTER	109	MKHMAGAAAAAGAVVGG	LGGYMLGS	AMSRPMMHF	NDWE	DR	YYR	ENMNRYPNQVYYR PVDQ
BOVINE	180	YSNQNNFVHDCVNI	TKQHTVTTTT	KG	ENFTET	DI	KMMER	VVEQMCITQYQRESQAYYCG
MAN	169	YSNQNNFVHDCVNI	TKQHTVTTTT	KG	ENFTET	DI	VKMMER	VVEQMCITQYQRESQAYYCG
SHEEP	172	YSNQNNFVHDCVNI	TKQHTVTTTT	KG	ENFTET	DI	KIMERV	VVEQMCITQYQRESQAYYCG
MOUSE	168	YSNQNNFVHDCVNI	TKQHTVTTTT	KG	ENFTET	DI	VKMMER	VVEQMCITQYQRESQAYYEG
HAMSTER	169	YNNQNNFVHDCVNI	TKQHTVTTTT	KG	ENFTET	DI	KIMERV	VVEQMCITQYQRESQAYYEG
BOVINE	239	- RGASVILFSSPPVILL	LISFLIFLIVG					
MAN	228	- RGSSMVLFS	SPPVILLISFLIFLIVG					
SHEEP	231	- RGASVILFSSPPVILL	LISFLIFLIVG					
MOUSE	228	- RRSSTVLFS	SPPVILLISFLIFLIVG					
HAMSTER	229	- RRSASVLFSSPPVILL	LISFLIFLIVG					

A 3D molecular model of a protein structure, likely a viral capsid, shown in a dark, textured surface. Three arrows point to specific regions: one labeled 'YYX' at the top, and two labeled 'YYR' on the left and right sides. A thin, light-colored line is drawn across the structure, possibly representing a specific pathway or interaction.

Figure 3

Solvent-accessible



Solvent-inaccessible

Figure 4

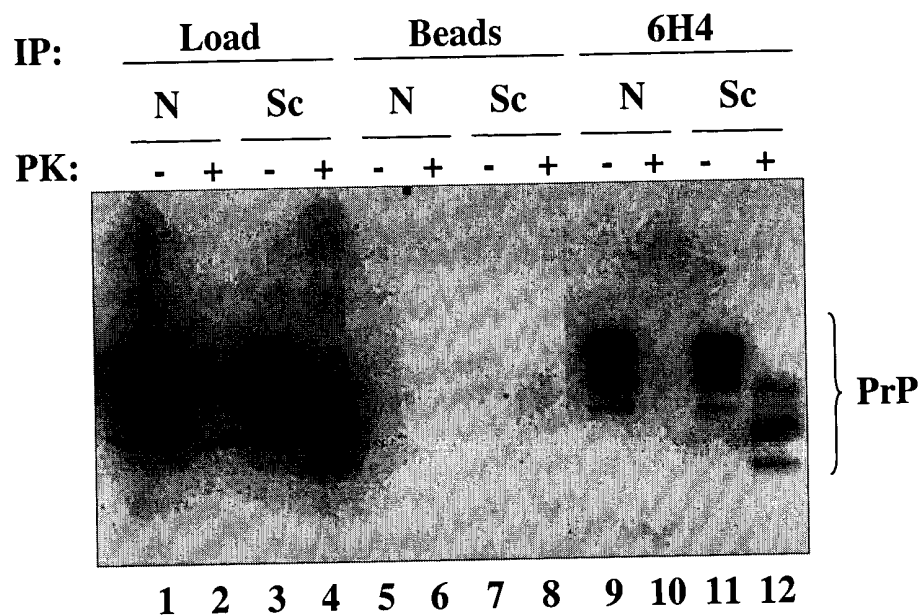


Figure 5

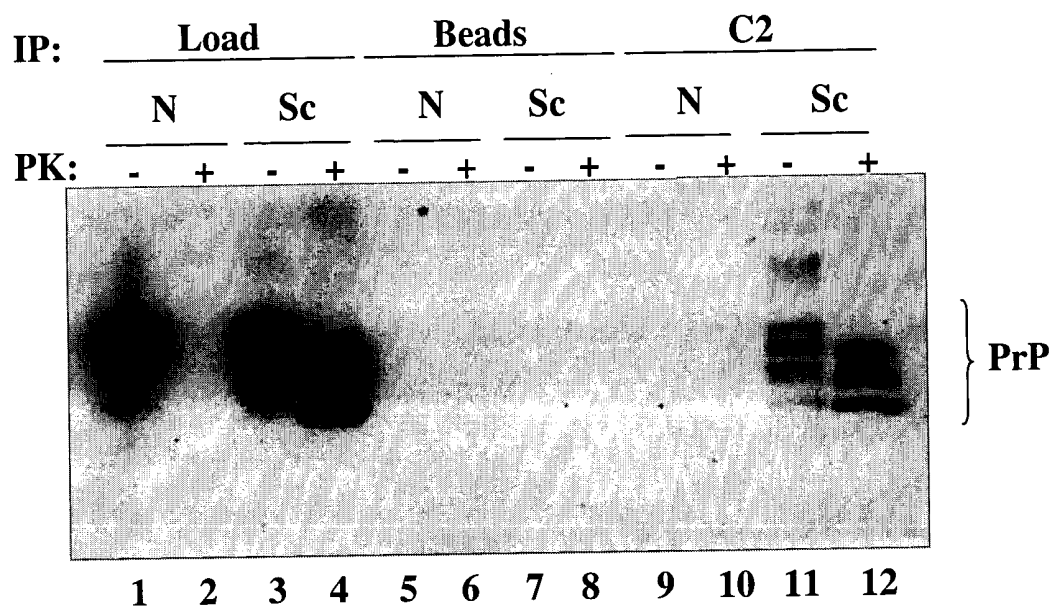


Figure 6

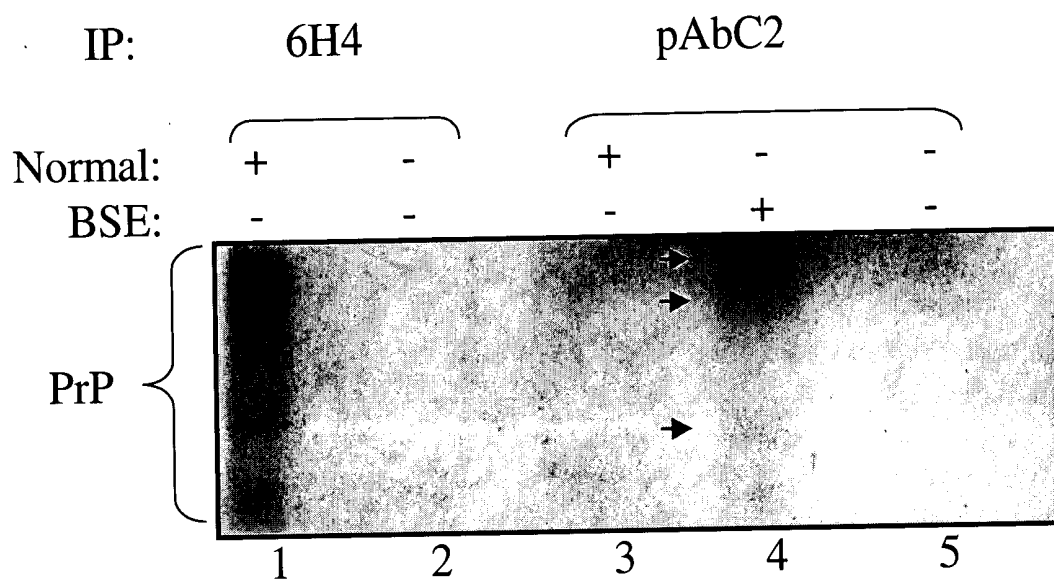


Figure 7

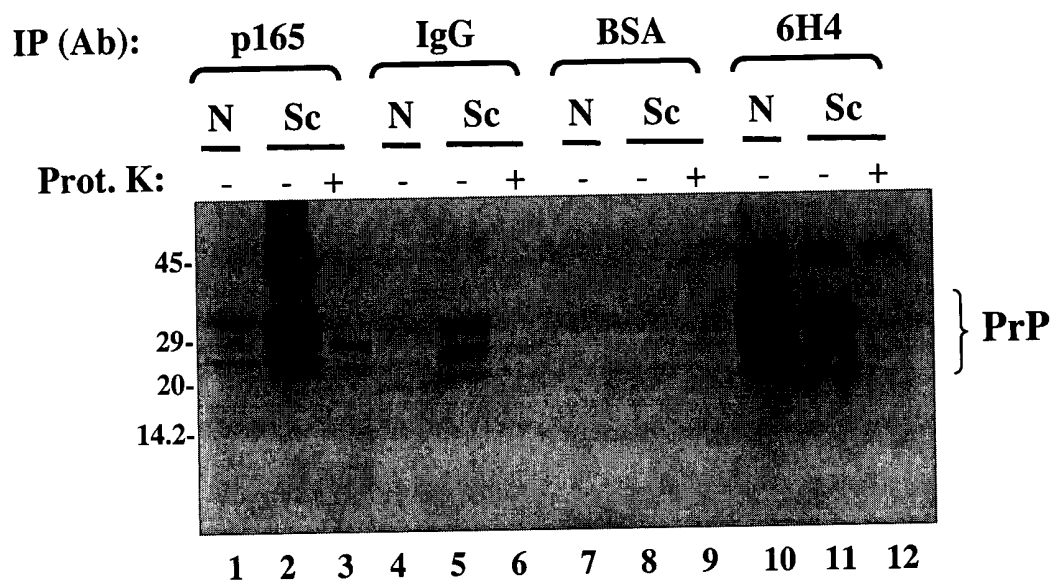


Figure 8

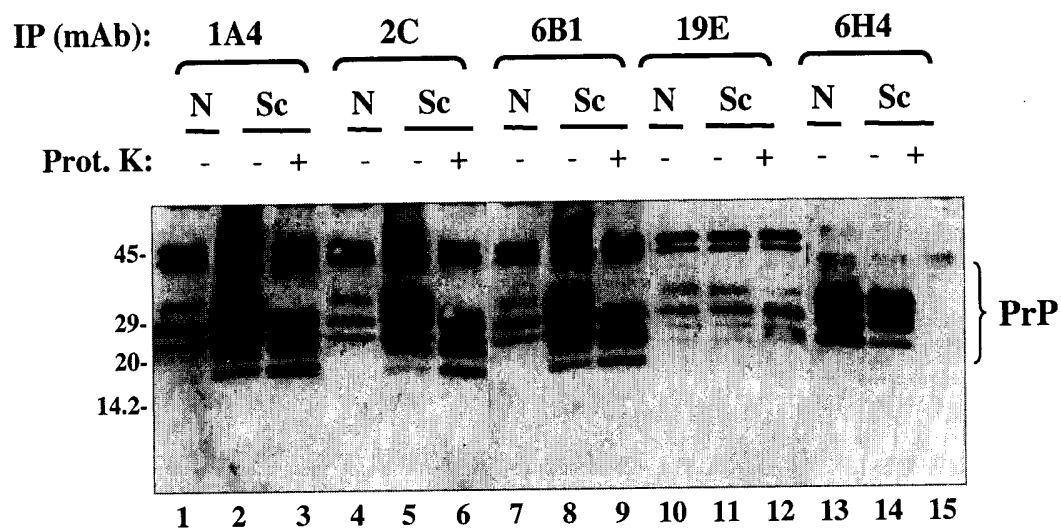
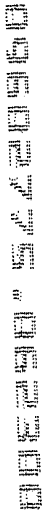


Figure 9

[illegible][illegible]

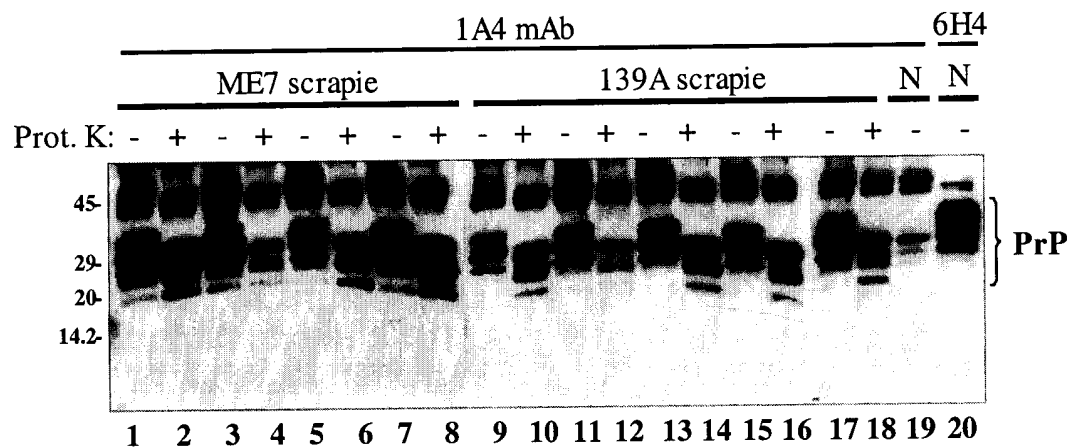
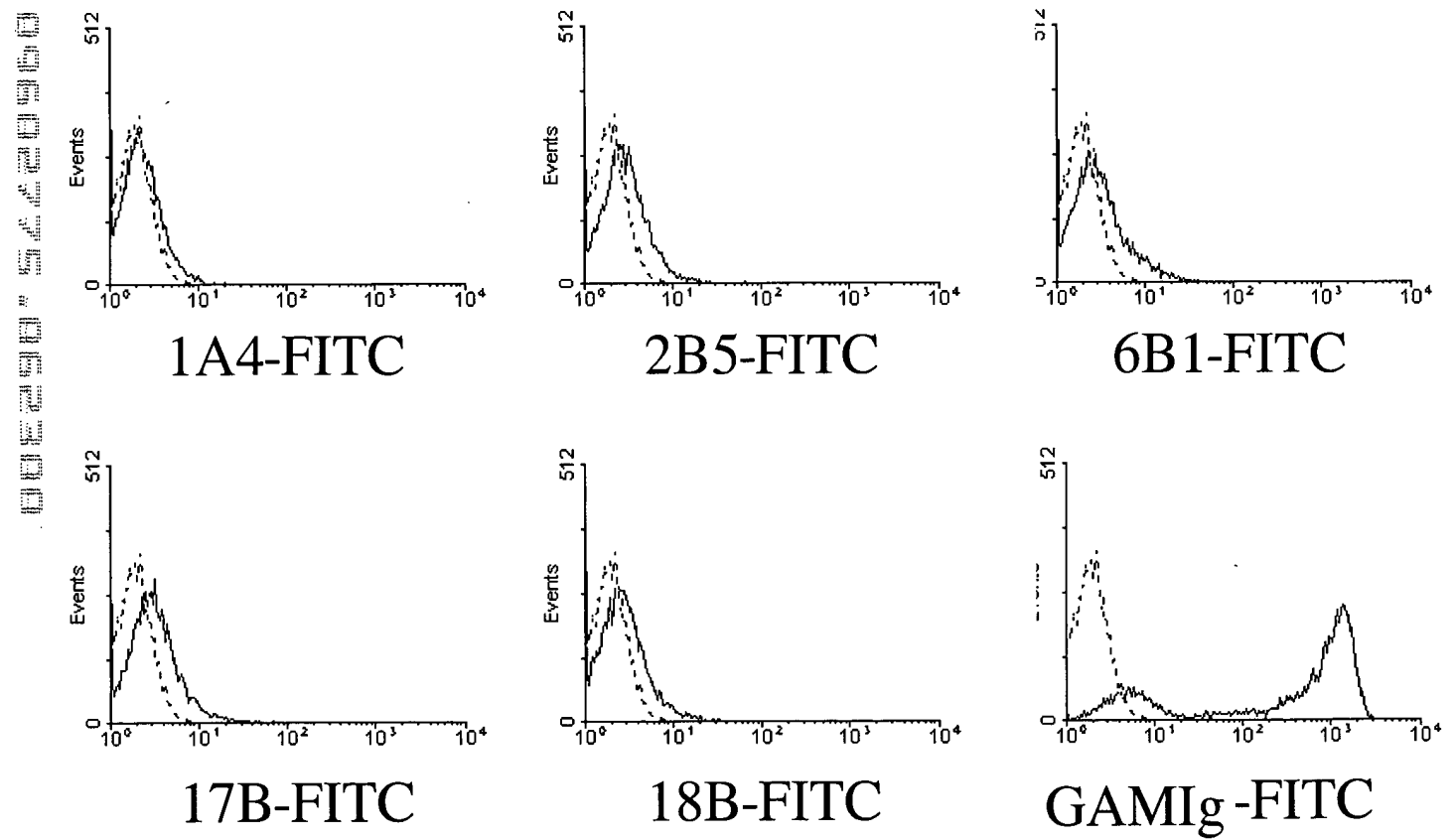


Figure 11

Figure 13



Western blot analysis of PrP in 1A4 and p165 cell lines. The blot shows protein bands for 1A4 (lanes 1-4) and p165 (lanes 5-10). Molecular weight markers are indicated on the left: 42 kDa, 30 kDa, 22 kDa, and 17 kDa. The right side of the blot is labeled 'PrP'. The lanes are grouped by cell line (1A4 and p165) and treatment (N and Sc). The 'Prot. K' treatment is indicated by '-' and '+' signs above the lanes. In the 1A4 panel, lanes 1 and 2 (N) show no band, while lanes 3 and 4 (Sc) show a strong band around 30 kDa. In the p165 panel, lanes 5 and 6 (N) show no band, while lanes 7 and 8 (Sc) show a strong band around 30 kDa. Lanes 9 and 10 (Sc) show a strong band around 30 kDa.

Figure 14

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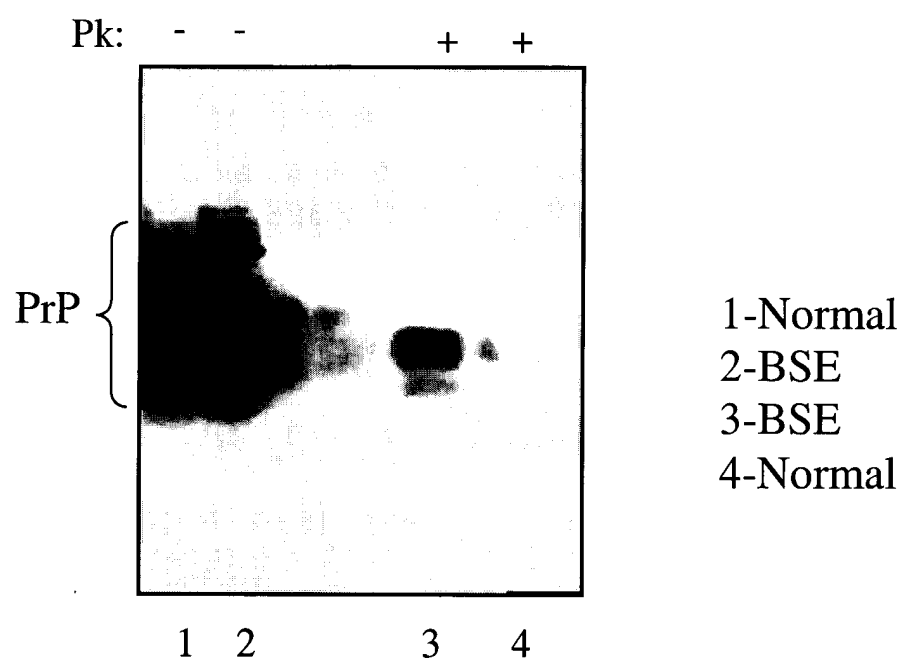


Figure 15

Figure 16

